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The *Caenorhabditis elegans* APC-related gene *apr-1* is required for epithelial cell migration and *Hox* gene expression

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Inactivation of the *Caenorhabditis elegans* APC-related gene (*apr-1*) has pointed at two separate functions of *apr-1*. First, *apr-1* is required for the migration of epithelial cells during morphogenesis of the embryo. In this process, APR-1 may act in a Cadherin/ α -Catenin/ β -Catenin complex as a component of adherens junctions. Second, *apr-1* is required for *Hox* gene expression, most likely by positively regulating the activity of the Wingless signaling pathway. During embryogenesis, *apr-1* is required for the expression of *ceh-13 labial* in anterior seam and muscle cells and during larval development, *apr-1* is necessary for the expression of *lin-39 deformed* in the vulval precursor cells. Thus, APR-1 may positively regulate the activity of the β -Catenin/Armadillo-related proteins HMP-2 in migrating epithelial cells and BAR-1 in the vulval precursor cells.

[Key Words: *Caenorhabditis elegans*; cell migration; vulva; APC; *Hox* gene; Wingless]

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The human APC tumor suppressor gene is mutated in most cases of inherited colorectal cancer (for review, see Kinzler and Vogelstein 1996; Polakis 1997). Humans carrying a germ-line mutation in one APC allele develop hundreds of colonic polyps at a young age, in which the second, wild-type APC allele has been lost spontaneously. Some of these polyps eventually accumulate additional mutations and progress to carcinomas. Spontaneous mutations in both APC alleles have also been observed in the majority of sporadic colon carcinomas. Biochemical studies have identified a number of APC-binding partners and suggested possible functions for APC. In particular, binding of APC to β -Catenin (Rubinfeld et al. 1993) is thought to regulate the activity of β -Catenin in the evolutionarily conserved Wingless (Wnt) signal transduction pathway (for review, see Polakis 1997; Wodarz and Nusse 1998; Peifer 1999). In both familial and sporadic forms of colorectal cancer, the vast majority of APC mutations that have been characterized result in carboxy-terminal truncations of the APC protein and eliminate part of the β -Catenin-binding site (Polakis 1995). Colon cancer cells that express truncated APC often exhibit elevated levels of cytoplasmic β -Catenin and an increased activity of the Wnt signaling pathway, suggesting that wild-type APC may negatively regulate β -Catenin function (Munemitsu et al. 1995; Ko-

rinek et al. 1997; Morin et al. 1997). Supporting this model, mutations in the *Drosophila* APC homologs dAPC1 and dAPC2 (or E-APC) result in the constitutive activation of the β -catenin homolog *armadillo* in different tissues (Ahmed et al. 1998; McCartney et al. 1999; Yu et al. 1999). However, studies of *Xenopus laevis* axis induction and early *Caenorhabditis elegans* embryonic development have suggested an activating rather than an inhibitory function for APC in the Wnt signaling pathway (Rocheleau et al. 1997; Vleminckx et al. 1997). In addition to its role in Wnt signaling, it has been proposed that vertebrate APC may control the migration and adhesion of epithelial cells (Nathke et al. 1996; Wong et al. 1996; Pollack et al. 1997), cell cycle progression (Baeg et al. 1995), and cell death (Morin et al. 1996).

During *C. elegans* development, activation of the Wnt signaling pathway induces the expression of *Hox* genes in different cell types (Eisenmann et al. 1998; Jiang and Sternberg 1998; Maloof et al. 1999). The *C. elegans* *Hox* genes are organized in an evolutionarily conserved gene cluster consisting of the four genes *ceh-13*, *lin-39*, *mab-5*, and *egl-5* that are the homologs of the *Drosophila* *Hox* genes *labial*, *deformed*, *antennapedia*, and *abdominal-B*, respectively (for review, see McGinnis and Krumlauf 1992; Lawrence and Morata 1994; Salser and Kenyon 1994). They are expressed in distinct domains along the anterior-posterior (A/P) axis and specify cell fates in their expression domains by controlling diverse processes such as cell division, polarity, migration, adhesion, fusion, and death. The β -catenin/armadillo-related

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gene *bar-1* is an essential regulator of the Wnt signaling pathway, leading to the induction of *Hox* gene expression during postembryonic development (Eisenmann et al. 1998; Maloof et al. 1999). In particular, *bar-1* is required for the transduction of a Wnt signal that specifies the vulval equivalence group in the six hypodermal Pn.p cells P3.p through P8.p (P3–8.p). In response to this Wnt signal, P3–8.p express the *Hox* gene *lin-39* and become vulval precursor cells (VPCs). The anterior Pn.p cells (P1.p and P2.p) and posterior Pn.p cells (P9.p–P11.p) that may not receive a Wnt signal do not express *lin-39* and adopt a fused (F) cell fate; they dissolve their adherens junctions and fuse with the surrounding hypodermal syncytium *hyp7* (Sulston and Horvitz 1977). In animals carrying a loss-of-function mutation in *bar-1*, P3–8.p frequently fail to express *lin-39* and adopt an F fate instead of a VPC fate (Eisenmann et al. 1998).

The *hmp-2* gene encodes another *C. elegans* β -Catenin homolog that is required for epithelial cell migration and elongation during morphogenesis of the embryo (Costa et al. 1998). HMP-2 does not appear to act in a Wnt signaling pathway, but rather, it functions as a component of the Cadherin/ α -Catenin/ β -Catenin complex at the adherens junctions of epithelial cells in the epidermis (which is called hypodermis in *C. elegans*).

We have taken a genetic approach to study the role of the *C. elegans* APC-related gene *apr-1* (Rocheleau et al. 1997) during embryogenesis and vulval development. Unlike the truncations found in human (Polakis 1995), mouse (Su et al. 1992), and *Drosophila* APC (Ahmed et al. 1998), the *apr-1* mutation described in this work is predicted to disrupt the synthesis of the entire APR-1 protein. Our analysis of the *apr-1* loss-of-function phenotype points at two separate functions of *apr-1*. First, *apr-1* is required for the migration and elongation of hypodermal cells that enclose the embryo during morphogenesis. In this context, APR-1 appears to function as a component of the hypodermal adherens junctions together with HMR-1 Cadherin, HMP-1 α -Catenin, and HMP-2 β -Catenin. Second, *apr-1* controls the expression of *Hox* genes during embryogenesis and vulval development, most likely by positively regulating the activity of the Wnt signaling pathway. In both processes, APR-1 may be necessary for the activity of β -Catenin/Armadillo-related proteins.

Results

Isolation of an *apr-1* loss-of-function mutation

A database search of the complete *C. elegans* genome sequence revealed the presence of a single APC homolog that had been previously named *apr-1* (Rocheleau et al. 1997). APR-1 and human APC1 both contain seven Armadillo repeats in their amino-terminal domains that are 31% identical and a conserved PDZ-binding motif at their carboxyl termini. The putative β -Catenin/Armadillo-binding site in the central region of APR-1 shows weaker sequence similarity when compared with human APC1.

To isolate a loss-of-function mutation in *apr-1*, a library consisting of $\sim 5 \times 10^5$ mutagenized animals was screened for deletions in the *apr-1* genomic region using a nested PCR assay (see Materials and Methods). A single allele, *apr-1(zh10)*, was isolated from this screen and used for the subsequent experiments. *apr-1(zh10)* animals contain a deletion of 1414 bp that removes the first, second, and most of the third exon (Fig. 1A). The *apr-1(zh10)* deletion is likely to completely eliminate gene function, as it removes the ATG translational initiation codon as well as the part of the 5' promoter region that contains a likely transcriptional start site with a TATA-box (Bensimhon et al. 1983).

The following observations indicate that loss of *apr-1* function causes a zygotic lethal phenotype. The progeny of *apr-1(zh10)/+* heterozygous parents contained no viable *apr-1(zh10)* homozygous animals as determined by PCR analysis ($n = 48$) or by scoring the segregation of a linked *unc-29(e1072)* mutation (see Materials and Methods). The lethality associated with *apr-1(zh10)* could be rescued by germ-line transformation with a plasmid containing a 7.3-kb genomic fragment of the *apr-1* locus (Fig. 1A). Furthermore, *apr-1(zh10)* homozygous animals only survived to adulthood if they carried an extrachromosomal array with the rescuing *apr-1* transgene.

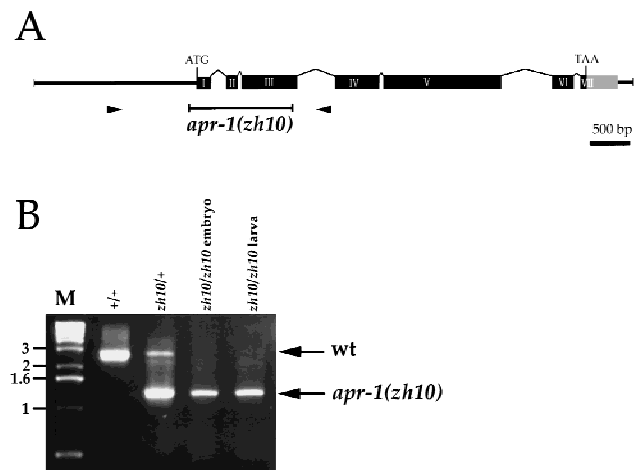


Figure 1. Genomic structure of the *apr-1* locus. (A) A 7.3-kb fragment (*apr-1wt*) containing all the sequences necessary to rescue the *apr-1* loss-of-function phenotype is shown. Exons are indicated by thick lines and are numbered. The positions of the ATG initiation and TAA stop codons are shown. Exon-intron boundaries were confirmed by RT-PCR analysis (data not shown). The extent of the 1414-bp deletion in *apr-1(zh10)* animals is indicated by the line underneath the genomic structure. The arrowheads indicate the position of the primers O95 and O101 that were used for the PCR analysis shown below. (B) Single-animal PCR analysis of a wild-type larva (+/+), a heterozygous *apr-1(zh10) unc-29(e1072)/lin-10(e1439)* larva (*zh10/+*), an *apr-1(zh10) unc-29(e1072)*-arrested embryo (*zh10/zh10* embryo) and an *apr-1(zh10) unc-29(e1072)*-arrested L1 larva (*zh10/zh10* larva). (M) Molecular weight standards; the numbers to the left of this lane indicate the length of relevant markers in kilobase pairs (kb). The arrows at right indicate the 2.9-kb product from the wild-type and the 1.5-kb product from the *zh10* allele.

To determine at what stage of development *apr-1(zh10)* mutants die, we monitored the development of a known number of eggs laid by *apr-1(zh10)/+* heterozygous parents. A total of $77 \pm 4\%$ of the eggs hatched and developed into viable animals of the genotype *apr-1(zh10)/+* or *+/+*, $15 \pm 3\%$ arrested during embryogenesis and $8 \pm 3\%$ arrested shortly after hatching during the first larval (L1) stage ($n = 450$). All arrested embryos and L1 larvae were homozygous for *apr-1(zh10)* as determined by PCR analysis ($n = 28$, Fig. 1B). Thus, approximately two-thirds of homozygous *apr-1(zh10)* mutants arrest during embryogenesis and the remaining third during the L1 stage. To test whether *apr-1(zh10)* L1 larvae may have escaped from the embryonic lethality because of rescue by maternal *apr-1* activity, we examined *apr-1(zh10)* germ-line mosaic animals (see Material and Methods). However, because loss of *apr-1* function in the germ line causes a sterile phenotype, we were unable to examine the contribution of maternal *apr-1* activity to embryogenesis.

Loss of *apr-1* function causes multiple morphogenesis defects in the embryo

During wild-type morphogenesis, contralateral pairs of hypodermal cells that are born on the dorsal side migrate ventrally along the left and right sides of the embryo until they reach the ventral midline and the hypodermis fully encloses the embryo, a process called ventral closure (Priess and Hirsh 1986; Williams Masson et al. 1997; Costa et al. 1998). At the same time, dorsal hypodermal cells intercalate and then fuse to each other, thereby forming the dorsal syncytium (Podbilewicz and White 1994; Williams Masson et al. 1998). After the cell migrations and fusions are completed, the hypodermis is organized in two rows of ventral and lateral epithelial cells on each side of the embryo plus a dorsal syncytium. Finally, the cells begin to elongate along the A/P axis until the length of the body has increased about fourfold.

apr-1(zh10) embryos that arrested before undergoing elongation ruptured on the ventral side, suggesting that *apr-1* may be required for ventral closure (Fig. 2A,B). To study the function of *apr-1* in hypodermal cells, *apr-1(zh10)* embryos were fixed at various stages and stained with the mAb MH27 that recognizes the adherens junction marker JAM-1 (Fig. 2C,D; Francis and Waterston 1985; M. Köppen and J.D. Hardin, pers. comm.). Alternatively, embryos carrying a *jam-1::GFP* reporter on the chromosomally integrated array *jcls1* were observed by four-dimensional time-lapse microscopy (Fig. 3; Mohler et al. 1998). Four kinds of defects could be observed in *apr-1(zh10)* embryos.

First, the migration of hypodermal cells was disrupted. Of 22 *apr-1(zh10)* embryos observed by four-dimensional microscopy, 18 arrested before undergoing elongation. In 13 of these 18 embryos, hypodermal cells initiated ventral migration, but they did not reach the ventral midline and failed to enclose the embryo (Fig. 3A,B). In the other five arrested embryos, the two pairs of anterior leading cells did reach the ventral midline and ventral closure

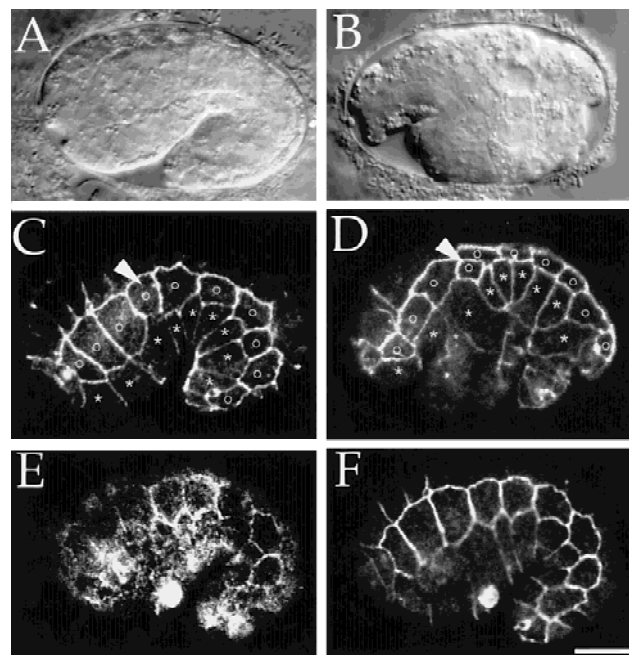


Figure 2. Embryonic phenotypes caused by the *apr-1(zh10)* deletion and APR-1 expression pattern in hypodermal cells. Nomarski photomicrographs of a wild-type embryo at the 1.5-fold stage (A) and a terminal stage *apr-1(zh10)* embryo that had ruptured ventrally and failed to elongate (B). Morphology of hypodermal cells stained with the mAb MH27 in a wild-type 1.5-fold embryo (C) and a terminal stage *apr-1(zh10)* embryo (D). The circles in C and D denote the lateral seam cells, the asterisks indicate the ventral hypodermal cells, and the arrowheads point at the position of the V1 seam cell. Note the rupture at the ventral midline, the dorsal retraction of ventral hypodermal cells, and the mislocalization of V1 (arrowhead) in the *apr-1(zh10)* embryo shown in D. Partial localization of APR-1 at the adherens junctions in a *zhEx2[apr-1wt]* embryo stained with APR-1 antibodies (E) and mAb MH27 (F). All embryos are shown from a lateral view; anterior is left and dorsal is up. The *apr-1(zh10)* embryos were the descendants of *apr-1(zh10) unc-29(e1072)/lin-10(e1439)* mothers. Scale bar in F, 10 μ m.

was initiated, but the embryos ruptured on the anterior or posterior ventral side at about the 1.25-fold stage. In all 18 terminal stage *apr-1(zh10)* embryos, the hypodermis had retracted dorsally and inner cells from the gut and other tissue protruded (Fig. 2D). The remaining four *apr-1(zh10)* embryos that enclosed and elongated may have hatched and arrested as L1 larvae as described above.

Second, in 7 of 10 *apr-1(zh10)* embryos, the dorsal pairs of hypodermal cells failed to intercalate (Fig. 3C,D; for the remaining 8 embryos, dorsal intercalation could not be scored because they were observed from a ventral view.). Although some cell fusions between the misaligned hypodermal cells occurred (arrowheads in Fig. 3C,D), no dorsal syncytium was formed in these seven embryos. All embryos exhibiting dorsal intercalation defects also failed to undergo ventral closure.

Third, hypodermal cells did not elongate along the A/P axis. At a time point when wild-type embryos had

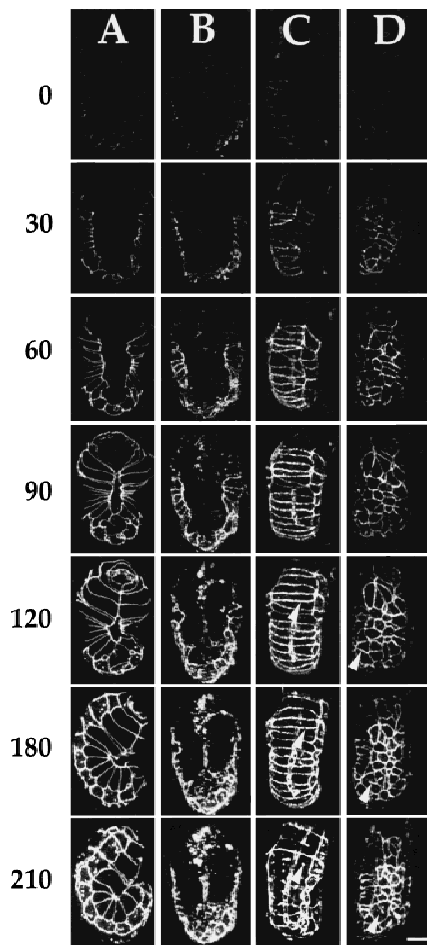


Figure 3. Ventral closure and dorsal intercalation defects in *apr-1(zh10)* embryos observed by four-dimensional microscopy. The migration of hypodermal cells in embryos carrying the adherens junction marker *jcIs1[jam-1::GFP]* was observed by two-photon excitation microscopy and four-dimensional reconstructions were done as described (Mohler and White 1998; Mohler et al. 1998). Ventral views of a developing wild-type (A) and an *apr-1(zh10)* embryo (B); dorsal views of a developing wild-type (C) and another *apr-1(zh10)* embryo (D). The time point when JAM-1::GFP could first be detected was taken as reference point 0 and is shown in the top frames of A–D. The numbers at left of the following frames indicate the time points relative to the reference point in minutes. Note in B the failure of the ventral hypodermal cells to reach the ventral midline and in D the lack of intercalation between the dorsal hypodermal cells. The arrowheads in C and D indicate fusing cells. The embryos were descendants of *apr-1(zh10) unc-29(e1072)/lin-10(e1439); jcIs1[jam-1::GFP]* mothers. In all embryos, anterior is up. Scale bar in D, 10 μ m.

elongated about two- to threefold, hypodermal cells in arrested *apr-1(zh10)* embryos had retained their square shape and the length of the embryos had not increased >1.25-fold (Fig. 3; data not shown).

Fourth, the lateral anterior hypodermis was frequently disorganized in *apr-1(zh10)* mutants. In wild-type embryos, 10 hypodermal cells (H0, H1, H2, V1–V6, and T) form the lateral row of seam cells on each side of the

embryo (Fig. 2C; Sulston et al. 1983). In 81% of *apr-1(zh10)* embryos, the two anterior seam cells H2 and V1 were mislocalized ($n = 47$). V1 was often shifted dorsally or ventrally (arrowhead in Fig. 2D) and H2 moved posterior to make direct contact with V2.

Similar ventral closure and elongation defects have been observed in *hmr-1* mutant embryos and in *hmp-1* or *hmp-2* mutants that lack zygotic and maternal gene activity (Costa et al. 1998). On the other hand, *hmr-1*, *hmp-1*, or *hmp-2* mutants do not exhibit dorsal intercalation defects or the misalignment of the anterior seam cells H2 and V1 that was observed in *apr-1(zh10)* mutants (Costa et al. 1998; our observation), suggesting that *apr-1* may perform additional functions in the dorsal hypodermis and in anterior seam cells that do not require *hmr-1*, *hmp-1*, or *hmp-2* activity. *hmr-1*, *hmp-1*, and *hmp-2* encode proteins that are similar to vertebrate Cadherins, α -Catenin, and β -Catenin, respectively, and are localized at hypodermal adherens junctions. We therefore asked whether APR-1 may also be a component of the hypodermal adherens junctions. Because the abundance of APR-1 was too low to allow the detection of endogenous APR-1 in immunohistochemistry experiments, multiple copies of wild-type *apr-1* were introduced on an extra-chromosomal array (*zhEx2*) to increase APR-1 expression levels. In *zhEx2* embryos, APR-1 was partially localized at adherens junctions, but some cytoplasmic staining could be detected as well (Fig. 2E,F). No staining could be observed with APR-1 preimmuneserum and the addition of an excess of soluble APR-1 antigen completely blocked the staining shown in Figure 2E.

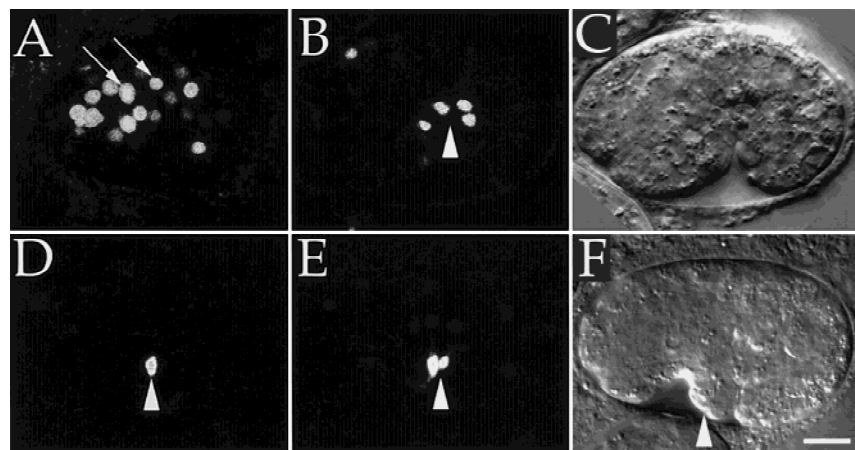
In summary, *apr-1* is necessary for at least four different processes during morphogenesis of the embryo, for the ventral migration, dorsal intercalation, and elongation of hypodermal cells and for the alignment of the anterior seam cells H2 and V1.

apr-1 is required for expression of the Hox gene ceh-13 during embryogenesis

A mutation in the *Hox* gene *ceh-13* causes a similar mislocalization of the anterior seam cells H2 and V1 (Brunschwig et al. 1999). *ceh-13* specifies cell fates in the anterior domain of the embryo, but it is not required for ventral closure, dorsal intercalation, or elongation of the embryo. Thus, one possible explanation for the mislocalization of the anterior seam cells H2 and V1 in *apr-1(zh10)* embryos is that *apr-1* may control the expression of the *Hox* gene *ceh-13*.

To test this hypothesis, we examined the expression pattern of CEH-13 in *apr-1(zh10)* embryos. We used a CEH-13::GFP reporter transgene on the chromosomally integrated array *swIs1* to observe CEH-13 expression (Wittmann et al. 1997). *swIs1[ceh-13::GFP]* embryos at the 1.25-fold stage showed strong GFP expression in the seam cells H2 and V1, in anterior muscle cells, and in ventral nerve cord (VNC) precursors in the central body region (Fig. 4A–C), similar to the expression pattern observed in embryos stained with CEH-13 antibodies (Brunschwig et al. 1999). In 82% of *apr-1(zh10)*;

Figure 4. *apr-1* controls *ceh-13* expression in anterior cells of the embryo. CEH-13::GFP expression in lateral (A) and middle (B) confocal sections through a wild-type 1.25-fold embryo and a Nomarski image of the same embryo (C). Lateral (D) and middle (E) confocal sections through an *apr-1(zh10)* embryo arrested before elongation and a Nomarski image of the same embryo (F). The arrows in A indicate the nuclei of H2 and V1. The arrowheads in B, D, and E point at the VNC precursors and in F at the site of ventral rupture. Note the absence of CEH-13::GFP expression in the lateral sections of the *apr-1(zh10)* embryo shown in D. Both embryos are shown from a lateral view; anterior is left and dorsal is up. The full genotype of the embryo shown in A–C was *him-5(e1490); swIs1[ceh-13::GFP]*; the *apr-1(zh10)* embryo shown in D–F was a descendant of an *apr-1(zh10) unc-29(e1072)/unc-13(e51) lin-11(n566); him-5(e1490); swIs1[ceh-13::GFP]* mother. Scale bar in F, 10 μ m.



swIs1[ceh-13::GFP] embryos, GFP expression was absent in anterior muscle and seam cells, whereas the VNC precursors continued to express CEH-13::GFP in all cases examined ($n = 39$, Fig. 4D–F). The loss of CEH-13 expression in the anterior seam cells H2 and V1 is unlikely to be due to the absence of these cells, because the expression pattern of the adherens junction marker JAM-1 indicated that H2 and V1 were always present in the anterior domain of *apr-1(zh10)* embryos (Fig. 2D). Thus, *apr-1* is necessary for expression of the *Hox* gene *ceh-13* in anterior cells but not in the VNC precursors that are located in the central body region.

APR-1 is expressed in the vulval precursor cells

During the early L1 stage, 6 pairs of hypodermal cells migrate along the right and left sides of the larva to the ventral midline, where they interdigitate and align in a row of 12 Pn cells along the A/P axis (P1 designates the anterior-most and P12 the posterior-most cell). Shortly after reaching the ventral midline, each Pn cell divides into a Pn.a neuroblast and a hypodermal Pn.p cell (Sulston and Horvitz 1977). Finally, a Wnt signal specifies the VPC fate in P3–8.p (Eisenmann et al. 1998).

The larval expression pattern of APR-1 was studied by staining animals carrying the *zhEx2[apr-1wt]* array with APR-1 antibodies or by observing GFP expression in a strain carrying an APR-1::GFP reporter transgene on the integrated array *zhIs2*. The adherens junctions were visualized by staining with the mAb MH27. In early L1 larvae, APR-1 was expressed in the 12 Pn cells as they were descending toward the ventral midline (Fig. 5A,B). In L3 stage hermaphrodites, APR-1 was strongly expressed in P3–8.p, but no APR-1 could be detected in anterior (P1.p and P2.p) or posterior (P9.p–P12.p) Pn.p cells (Fig. 5C,D; data not shown). In addition, APR-1 was expressed in the seam cells (arrowheads in Fig. 5C,D), in the excretory cell and in the excretory canal cell during the L1 stage (horizontal arrow in Fig. 5A,B), and in some unidentified neurons in the head region (not shown).

Similar to the staining observed in embryos, some APR-1 staining was localized at the adherens junctions, whereas another fraction of APR-1 could be detected in the cytoplasm (Fig. 5C–F). The specific expression of APR-1 in the VPCs raised the possibility that *apr-1* may act during vulval development.

apr-1 is necessary for the specification of the vulval equivalence group

We devised two different approaches to inactivate *apr-1* function in the Pn.p cells without affecting *apr-1* activity during embryogenesis. First, we used *cre/loxP*-mediated recombination to inactivate a rescuing *apr-1* transgene (Kilby et al. 1993). To this end, *loxP* sites were introduced into the rescuing *apr-1* construct such that *cre*-mediated recombination would result in the excision of all but the first exon of *apr-1* (*loxP* > *apr-1* > *loxP*; Fig. 6A). *cre* recombinase was expressed under control of the Pn.p cell-specific *lin-31* promoter (*lin-31::cre*; Tan et al. 1998). The *loxP* > *apr-1* > *loxP* and *lin-31::cre* plasmids were introduced into *apr-1(zh10)* animals on two separate extrachromosomal arrays, yielding a strain of the genotype *apr-1(zh10); zhEx11[loxP > apr-1 > loxP]; zhEx12[lin-31::cre]* (see Materials and Methods). In single animals from this strain, the expected recombination product could be detected by PCR analysis (Fig. 6A). As a second approach, we performed tissue-specific RNA-mediated interference (RNAi; Fire et al. 1991). For this purpose, a plasmid carrying anti-sense *apr-1* cDNA under control of the Pn.p cell-specific *lin-31* promoter (*lin-31::apr-1i*; Fig. 6B) was introduced into wild-type animals on the integrated array *galIs44*. No APR-1 protein was detectable in the VPCs of *galIs44; zhEx2[apr-1wt]* animals, whereas the seam cells in the same animals continued to express APR-1 as determined by antibody staining (data not shown).

Both approaches yielded animals displaying essentially the same vulval phenotype, although with a different penetrance. In 15% of *apr-1(zh10); zhEx11[loxP > apr-*

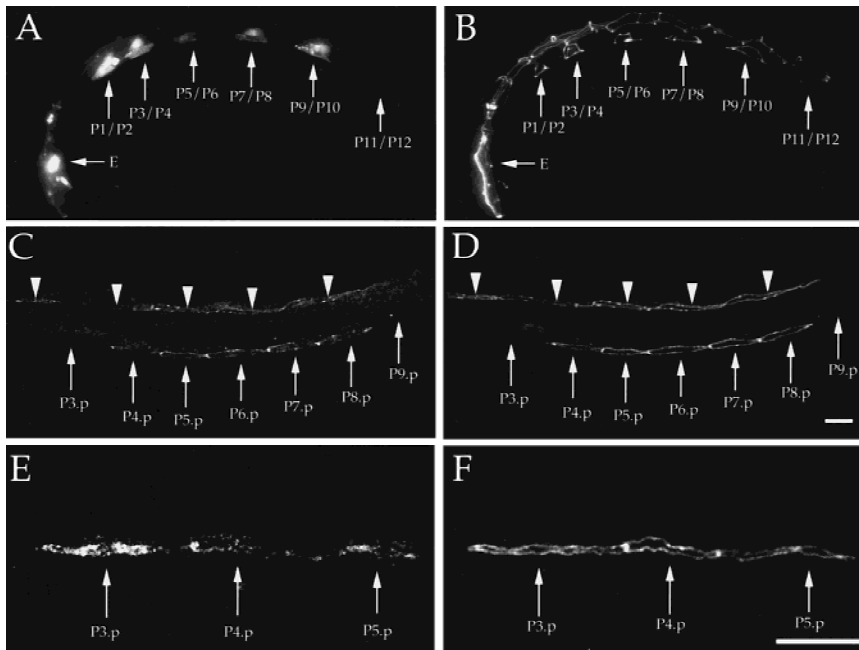


Figure 5. APR-1 expression pattern during larval development. (A) APR-1::GFP expression and (B) mAb MH27 staining of an L1 larva carrying the *zhEx2[apr-1::GFP]* array. The vertical arrows in A and B point at the six pairs of P cells that were descending from the *left* and *right* sides toward the ventral midline, the horizontal arrow points at the excretory cell body. (C) APR-1 antibody and (D) mAb MH27 staining of an L3 larva carrying the *zhEx2[apr-1wt]* array. Note the APR-1 expression in the VPCs (P4–8.p) indicated by the vertical arrows and in the seam cells indicated by the arrowheads. In this animal, P3.p had adopted an F fate (as observed in 39% of wild-type animals) and did not express APR-1. No APR-1 expression was detectable in the anterior (P1–2.p) or posterior (P9–11.p) Pn.p cells (P1.p, P10.p, and P11.p are not shown). (E) Higher magnification of APR-1 antibody, and (F) mAb MH27 staining of P3–5.p in an animal in which P3.p had adopted a VPC fate. All larvae are shown from a lateral view, anterior is *left*. Scale bars in D and F, 10 μ m.

1 > *loxP*]; *zhEx12[lin-31::cre]* and in 86% of *gals44[lin-31::apr-1i]* animals, P3–8.p adopted an F cell fate instead of induced (1° or 2°) or uninduced (3°) VPC fates (Sternberg and Horvitz 1986) as determined by staining of the adherens junctions with the mAb MH27 (Fig. 6C–E; Table 1) or by directly observing the pattern of cell divisions (Table 2, rows 2–6). Fusion of P3–8.p could be detected early in the L2 stage as opposed to the later fusion of P3.p that occurred in 39% of wild-type larvae at the beginning of the L3 stage (Table 1, row 1; Sternberg and Horvitz 1986; Eisenmann et al. 1998). Direct observation of the cell lineage of P3–8.p in *gals44[lin-31::apr-1i]* animals confirmed that cells that presumably adopted an F fate did not divide and remained at their original position in the ventral midline throughout vulval development (Table 2). With both approaches, Pn.p cells that were distal to the anchor cell (P3.p and P4.p) were more likely to adopt an F fate than those proximal to the anchor cell (P6.p, followed by P5.p, P7.p, and P8.p; Tables 1 and 2). From these experiments, we have concluded that *apr-1* acts in the Pn.p cells to specify the VPC fate in P3–8.p.

Beside *apr-1*, the *Hox* gene *lin-39* *deformed* and the β -catenin/*armadillo*-related gene *bar-1* are required for specification of the vulval equivalence group (Clark et al. 1993; Wang et al. 1993; Eisenmann et al. 1998). Because a null mutation in *bar-1* causes an incompletely penetrant phenotype, we asked whether inactivation of *apr-1* in *bar-1(0)* animals might increase the frequency of the F fate in P3–8.p. In *gals44[lin-31::apr-1i]; bar-1(0)* double mutants (Table 1, row 6), P3–8.p more frequently adopted an F fate than in either *bar-1(0)* or *gals44[lin-31::apr-1i]* single mutants (Table 1, rows 4,5), suggesting that in the absence of *bar-1* function, *apr-1* may still be necessary for the specification of the VPC fate.

apr-1 is required for the expression of the *Hox* gene *lin-39* in P3–8.p

Because *bar-1* regulates *lin-39* expression in response to a Wnt signal (Eisenmann et al. 1998), we tested whether *apr-1* may also be required to induce *lin-39* expression in P3–8.p. The expression pattern of LIN-39 was determined by staining L2 and L3 larvae with LIN-39 antibodies (Maloof and Kenyon 1998). In wild-type animals, P3–8.p expressed LIN-39, except for those L3 larvae in which P3.p had adopted an F fate (Fig. 7A,C,E). In all 5 *apr-1(zh10); zhEx11[loxP > apr-1 > loxP]; zhEx12[lin-31::cre]* and all 50 *gals44[lin-31::apr-1i]* animals, no LIN-39 expression was detectable in Pn.p cells that had adopted an F fate. An example is shown in Fig. 7, B and D. Consistent with the increasing frequency of the F fate in Pn.p cells distal to the anchor cell, LIN-39 was undetectable in P3.p but strongly expressed in P6.p in most *gals44[lin-31::apr-1i]* animals (Fig. 7F). Furthermore, the down-regulation of LIN-39 expression could be observed in late L1 larvae before Pn.p cells adopted an F fate (10–12 hr posthatching, data not shown), indicating that the loss of LIN-39 expression is a cause and not a consequence of the F fate. In summary, these data have suggested that *apr-1* may act in a Wnt signaling pathway that induces LIN-39 expression in P3–8.p.

Discussion

Our genetic analysis of the APC-related gene *apr-1* has revealed distinct functions of *apr-1*: First, *apr-1* is required for the ventral migration, dorsal intercalation, and elongation of hypodermal cells during morphogenesis of the embryo (Fig. 8A). Second, *apr-1* regulates *Hox*

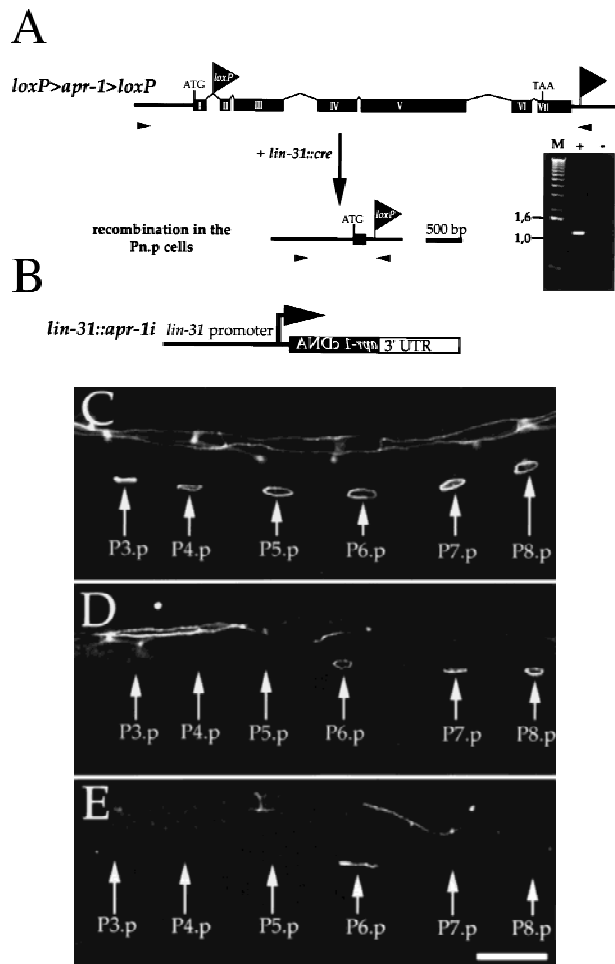


Figure 6. Inactivation of *apr-1* in the Pn.p cells causes P3–8.p to adopt an F fate. (A) Structure of the *loxP > apr-1 > loxP* transgene used for *cre/loxP*-mediated inactivation of *apr-1* and of the predicted recombination product. The circular excision product is not shown. The arrowhead below the *loxP > apr-1 > loxP* construct indicates the position of the primers O101 (see Materials and Methods) and M13reverse (Invitrogen) that were used for the PCR analysis shown at right. (+) The 1.3-kb recombination product obtained by PCR analysis of an L4 larva of the genotype *apr-1(zh10); zhEx11[loxP > apr-1 > loxP]; zhEx12[lin-31::cre]*; (–) an animal of the genotype *apr-1(zh10); zhEx11[loxP > apr-1 > loxP]* analyzed as negative control. The PCR conditions used did not allow amplification of the 6.5-kb product from *loxP > apr-1 > loxP* transgenes that had not undergone *cre*-mediated recombination. (M) Molecular mass standards; the numbers at left indicate the length of relevant markers in kilobase pairs. (B) Structure of the *lin-31::apr-1* construct used for Pn.p cell-specific RNAi. mAb MH27 staining of the VPCs in wild-type (C), *apr-1(zh10)*; *zhEx11[loxP > apr-1 > loxP]; zhEx12[lin-31::cre]* (D), and *gal4[lin-31::apr-1]* L2 larvae (E). The arrows point at the positions of the Pn.p nuclei as determined by parallel staining with DAPI (not shown). Note that in the animal shown in D P3.p, P4.p, and P5.p and in the animal shown in E, all Pn.p cells except for P6.p have adopted an F fate, because they failed to stain with the mAb MH27. See Table 1 for more examples and for the complete genotypes of the animals shown. Scale bar in E, 10 μ m.

gene expression. During embryogenesis, *apr-1* is required for the expression of the *Hox* gene *ceh-13 labial* in anterior cells (Fig. 8A) and during vulval development, *apr-1* is necessary for the expression of the *Hox* gene *lin-39 deformed* in P3–8.p (Fig. 8B).

apr-1 is required for morphogenesis of the embryo

apr-1 plays multiple roles during morphogenesis of the embryo. The incomplete penetrance of the different phenotypes is probably due to a variable rescue by maternally expressed *apr-1*. However, because loss of *apr-1* function in the germ line causes complete sterility, we were unable to examine the contribution of maternal *apr-1* activity to morphogenesis.

The ventral closure defect is caused by a disruption of hypodermal cell migration. In *apr-1(zh10)* embryos, hypodermal cells initiate ventral migration, but they stop before reaching the ventral midline. Moreover, in *apr-1(zh10)* larvae that arrest during the L1 stage the hypodermal Pn cells do not descend to the ventral midline, suggesting that *apr-1* may also be required for the ventral migration of the Pn cells (data not shown). What may be the role of *apr-1* in hypodermal cell migration? One possible model predicts that APR-1 may act in a Cadherin/ α -Catenin/ β -Catenin complex at the adherens junctions. It has been proposed that filopodial contacts must be made between the contralateral pairs of migrating cells before adherens junctions can be formed across the ventral midline and the cells are fixed in their new position (Costa et al. 1998; Raich et al. 1999). In particular, the two pairs of leading cells project actin-rich filopodia toward the ventral midline (Williams Masson et al. 1997). APR-1 may be required together with HMR-1 Cadherin, HMP-1 α -Catenin, and HMP-2 β -Catenin to stabilize the transient filopodial contacts between the pairs of leading cells by attaching actin filaments to the newly forming junctions.

Dorsal intercalation does not depend on *hmr-1*, *hmp-1*, or *hmp-2* activity, possibly because this process does not involve the formation of new adherens junctions. *apr-1*, on the other hand, may be required for the directed movement of dorsal hypodermal cells through a microtubule-dependent mechanism that has been proposed for vertebrate APC (Nathke et al. 1996; Pollack et al. 1997). In migrating vertebrate cells, APC was found in clusters near the ends of microtubules that extended into regions of actively migrating plasma membranes and at the sites where two cells first touched. This specific subcellular localization has suggested that APC may recruit microtubules into the extending membrane structures and thus stabilize the direction of cell migration. Although we have not observed a specific localization of APR-1 at the sites of actively migrating membranes, our data are consistent with an analogous role for *apr-1* in directing cell movements during ventral closure and dorsal intercalation.

An alternative model explaining both the ventral closure and dorsal intercalation defects predicts that *apr-1* may be required for the specification of hypodermal cell

Table 1. *Pn.p* cells lacking *apr-1* activity frequently adopt an F fate

| Row | Genotype | Percent fused ^a | | | | | | Percent mutant ^b | No. ^c |
|-----|---|----------------------------|------------------|------|------|------------------|------|-----------------------------|------------------|
| | | P3.p | P4.p | P5.p | P6.p | P7.p | P8.p | | |
| 1 | wild type | 39 | 0 | 0 | 0 | 0 | 0 | 0 | 61 ^d |
| 2 | <i>apr-1; zhEx11[loxP>apr-1>loxP]</i> ^e | 33 | 1.4 ^f | 0 | 0 | 0.7 ^f | 0 | 1.4 | 140 |
| 3 | <i>apr-1; zhEx11[loxP>apr-1>loxP]; zhEx12[lin-31::cre]</i> ^g | 45 | 9 | 4 | 3 | 4 | 4 | 15 | 219 |
| 4 | <i>gaIs44[lin-31::apr-li]</i> | 72 | 60 | 44 | 17 | 35 | 35 | 86 | 57 |
| 5 | <i>bar-1(0)</i> ^h | 66 | 56 | 11 | 2 | 2 | 6 | 62 | 55 |
| 6 | <i>bar-1(0); gaIs44[lin-31::apr-li]</i> ^h | 95 | 88 | 81 | 61 | 54 | 63 | 99 | 103 |

For each strain, L3 larvae were stained with mAb MH27 to detect the adherens junctions of unfused cells.

^aEach column indicates the percent frequency at which Pn.p cells adopted an F fate instead of a VPC fate.

^bThe frequency at which animals with fewer than six VPCs were observed in each strain; animals in which P3.p had adopted an F fate were counted as wild-type.

^cThe number of animals scored for each strain.

^dSimilar data for wild-type (N2) have been reported previously (Eisenmann et al. 1998).

^eThe complete genotype of this strain was *apr-1(zh10) unc-29(e1072); zhEx11[loxP>apr-1>loxP]*.

^fIn two animals of this strain, P4.p and/or P7.p had adopted an F fate. This could be due to a spontaneous loss of the *zhEx11* array in single Pn.p cells or to an incomplete rescue by the *apr-1* transgene. In other strains carrying a rescuing *apr-1* transgene, individual Pn.p cells adopted an F fate in 0.4%–0.8% of the animals (data not shown).

^gThe complete genotype of this strain was *apr-1(zh10) unc-29(e1072); unc-119(e2498); zhEx11[loxP>apr-1>loxP]; zhEx12[lin-31::cre, unc-119(+)]*.

^hThe *bar-1(ga80)* null mutation was used in these experiments (Eisenmann et al. 1998).

fates. In particular, *apr-1* may be necessary for the transduction of a Wnt signal that specifies hypodermal cell fates in the embryo (see below). The disruption of cell migration and intercalation could therefore be the consequence of cell fate transformations. This scenario seems most likely in the case of dorsal intercalation because the dorsal hypodermal cells in *apr-1* mutants appear to be disorganized even before intercalation normally begins.

The regulation of cell migration by APC is of particular importance in the human intestine, because defects in epithelial cell migration may result in the formation of intestinal polyps (Kinzler and Vogelstein 1996). At the base of the crypts in intestinal epithelia, stem cells continuously proliferate to generate enterocytes that migrate out of the crypts up to the adjacent villi, where they undergo apoptosis. Mutations in APC may lead to the formation of intestinal polyps because they prevent the migration of enterocytes from the proliferative zone at the base of the crypts to the apoptotic zone in the villi,

leading to the accumulation of enterocytes in the crypts (Nathke et al. 1996).

In addition to the ventral migration and dorsal intercalation defects, hypodermal cells fail to elongate in *apr-1* mutants. Elongation may only be possible after ventral closure and dorsal intercalation have been completed. Alternatively, APR-1 may be required together with HMR-1, HMP-1, and HMP-2 to attach actin filaments to the adherens junctions of the hypodermis. The contraction of these actin filaments along the dorsoventral axis generates the force elongating hypodermal cells along the A/P axis, thus driving elongation of the entire embryo (Priess and Hirsh 1986).

apr-1 is required for the expression of the Hox genes *ceh-13* and *lin-39*

Loss of *apr-1* function during embryogenesis causes defects in the organization of the anterior seam cells, similar to a mutation in the *Hox* gene *ceh-13* (Brunschwig et

Table 2. Vulval Lineages of Wild-type, *gaIs44[lin-31::apr-li]*, and *lin-39(0)* animals

| Row | Genotype | P3.p | P4.p | P5.p | P6.p | P7.p | P8.p | Phenotype |
|-----|-------------------------------|----------|------|------|------|------|------|------------------|
| 1 | wild type | S S or F | S S | LLTN | TTTT | NTLL | S S | non-Vul |
| 2 | <i>gaIs44[lin-31::apr-li]</i> | F | F | LLTN | TTTT | F | F | Vul ^a |
| 3 | <i>gaIs44[lin-31::apr-li]</i> | F | F | LLTN | OOOO | F | SOO | Vul |
| 4 | <i>gaIs44[lin-31::apr-li]</i> | F | F | F | TTTT | F | F | Vul |
| 5 | <i>gaIs44[lin-31::apr-li]</i> | F | F | F | OOOO | F | F | Vul |
| 6 | <i>gaIs44[lin-31::apr-li]</i> | F | F | F | F | F | F | Vul |
| 7 | <i>lin-39(0)</i> | F | F | F | F | F | F | Vul ^b |

The cleavage planes of the third round of cell divisions are indicated as follows: (T) Transverse division; (O) oblique division; (L) longitudinal division; (N) cell that did not divide during the third round of division; (S) cell that fused with hyp7 after the first round of cell division; (F) cell that never divided. The shaded area indicates cells that adopted induced (1°, 1°-like, or 2°) cell fates.

^aWhen inspected under a dissecting microscope, 80% of *gaIs44[lin-31::apr-li]* animals displayed a Vul phenotype ($n > 200$).

^bData from Clark et al. (1993).

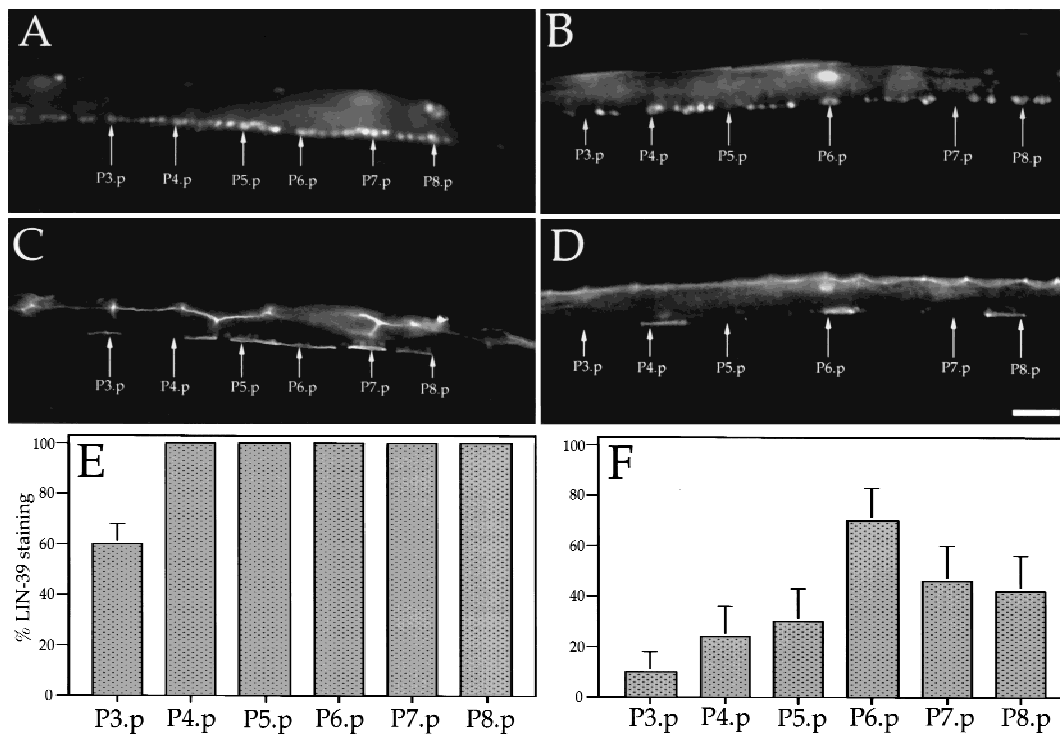


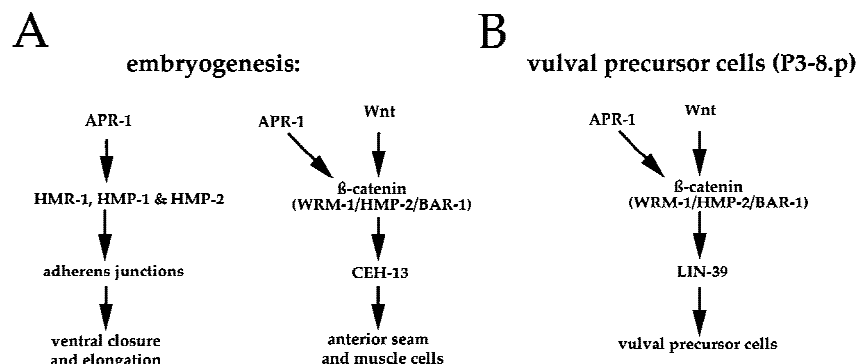
Figure 7. *apr-1* regulates *lin-39* expression in P3–8.p. LIN-39 antibody staining of wild-type (A) and *galIs44[lin-31::apr-1]* L2 larvae (B). The arrows point at the positions of the Pn.p nuclei as determined by parallel staining with DAPI (not shown). (C,D) mAb MH27 staining of the animals shown in A and B, respectively. Note that in the *galIs44[lin-31::apr-1]* animal shown in B and D, P3.p, P5.p, and P7.p failed to express LIN-39 and adopted an F fate as indicated by the absence of adherens junctions. LIN-39 expression in the individual Pn.p cells was scored in 50 wild-type (E) and 50 *galIs44[lin-31::apr-1]* L3 larvae (F) and the frequencies of LIN-39-positive cells are given as percent values. Error bars, the upper 95% confidence intervals. Scale bar in D, 10 μ m.

al. 1999). Inactivation of *apr-1* in the Pn.p cells causes P3–8.p to adopt an F cell fate, similar to mutations in the *Hox* gene *lin-39* (Clark et al. 1993; Wang et al. 1993). In both cases, *apr-1* is necessary for the expression of the respective *Hox* genes (Fig. 8A,B).

A Wnt signal that is transduced by the β -catenin-related gene *bar-1* induces expression of *lin-39* in P3–8.p (Eisenmann et al. 1998). Thus, *apr-1* may act in the Wnt signaling pathway in P3–8.p. Inactivation of *apr-1* in the

Pn.p cells by RNAi does not cause all six of the P3–8.p cells to adopt an F fate, suggesting that *apr-1* may not be absolutely required for Wnt signaling. Pn.p cells that are distal to the anchor cell more frequently adopt an F fate and down-regulate *lin-39* expression than those proximal to the anchor cell. A similar bias in the frequency of F fates has been observed in *bar-1* mutants (Eisenmann et al. 1998). One possible explanation for this finding is that proximal Pn.p cells may be closer to the source of the

Figure 8. Models for the different functions of APR-1. (A) During morphogenesis of the embryo, APR-1 may perform at least two separate functions. First, (left) APR-1 may act together with the adherens junction proteins HMR-1, Cadherin, HMP-1 α -Catenin, and HMP-2 β -Catenin in migratory hypodermal cells during ventral closure and elongation of the embryo. Second (right), APR-1 may act in the Wnt signaling pathway that controls the expression of the HOX protein CEH-13 in the anterior seam and muscle cells. (B) During larval development, APR-1 may be required for the transduction of a Wnt signal that induces expression of the HOX protein LIN-39 in P3–8.p to specify the VPC fate. The three β -Catensins WRM-1 (Rocheleau et al. 1997), HMP-2 (Costa et al. 1998), and BAR-1 (Eisenmann et al. 1998) may act redundantly in the transduction of Wnt signals.



Wnt signal in the central body region and thus require less *apr-1* or *bar-1* activity than distal Pn.p cells. Moreover, because loss of *bar-1* function results in an incompletely penetrant vulval phenotype, one of the other two β -catenin homologs (*hmp-2* or *wrm-1*) may be partially redundant with *bar-1* in the transduction of a Wnt signal in P3–8.p. Consistent with this hypothesis, inactivation of *apr-1* in animals carrying a *bar-1* loss-of-function mutation appears to further decrease the activity of the Wnt signaling pathway.

Less is known about a putative Wnt signal that may control *ceh-13* expression during embryogenesis. However, the *ceh-13* gene contains a putative LEF-1/Tcf-binding site that plays a role in the transcriptional activation of *ceh-13* in the E lineage (R. Köhler and F. Müller, pers. comm.). Because the Wnt signaling pathway controls the activity of the LEF-1/Tcf family of transcription factors (Brunner et al. 1997; van de Wetering et al. 1997), *ceh-13* expression may be regulated by Wnt signals. Furthermore, expression of the *ceh-13* homolog *labial* in the *Drosophila* midgut is controlled by a Wnt signal (Bienz 1994). These observations together with our data suggest that *apr-1* may be required to transduce a Wnt signal that induces *ceh-13* expression in anterior cells of the embryo (Fig. 8A). It is not clear which of the three *C. elegans* β -catenin/armadillo homologs may participate in the transduction of this putative Wnt signal. *hmp-2* mutant embryos do not exhibit defects in the alignment of seam cells (Costa et al. 1998) and they continue to express *ceh-13* until the terminal stage (our observation), *bar-1* mutant embryos do not exhibit an embryonic phenotype (Eisenmann et al. 1998) and no *wrm-1* mutation has been described to date (Rocheleau et al. 1997). However, two or all three of the β -catenin/armadillo-related genes might act redundantly.

APR-1 is necessary for β -Catenin/Armadillo function

Vertebrate and *Drosophila* APCs bind to β -Catenin/Armadillo proteins and promote their proteolytic degradation (Munemitsu et al. 1995; Polakis 1997). Mutations that remove the β -Catenin-binding site of APC cause the cytoplasmic accumulation of β -Catenin, resulting in the constitutive activation of the Wnt signaling pathway. Thus, APC is thought to antagonize β -Catenin/Armadillo function.

Our data and a prior analysis of maternal *apr-1* activity in early embryonic development by RNAi (Rocheleau et al. 1997) have indicated an opposite function for APR-1. In all cases that were studied, loss of *apr-1* function resulted in essentially the same phenotype as loss of β -catenin function. In particular, we have found that APR-1 is necessary for HMP-2 function during morphogenesis of the embryo and BAR-1 function in the transduction of a Wnt signal in P3–8.p.

How could this apparent reversal of APR-1 activity in comparison with APC in other species be explained? One possibility is that, although APR-1 is most similar to vertebrate and *Drosophila* APC, *apr-1* may not be a functional APC homolog. Because no second gene encod-

ing a protein with significant homology to APC has been found in the *C. elegans* genome, β -Catenins in *C. elegans* may not be subject to the same kind of negative regulation by APC as in other organisms. On the other hand, the activating function of APR-1 may reflect a unique requirement of the *C. elegans* β -Catenin homologs.

Another model predicts that APC may activate as well as inhibit β -Catenin/Armadillo function. The *apr-1* loss-of-function mutation described in this work may disrupt both activities, and loss of the activating function may prevail over loss of the inhibitory function. In contrast, the carboxy-terminal truncations found in vertebrate and *Drosophila* APC (Su et al. 1992; Polakis 1995; Ahmed et al. 1998) may selectively disrupt the inhibitory function of APC, resulting in the constitutive activation of β -Catenin/Armadillo. Supporting this hypothesis, overexpression of human APC in *Xenopus laevis* embryos induces axis duplication, as does overexpression of β -Catenin or Xwnt-8, suggesting that human APC may also have an activating function in the Wnt signaling pathway (Vlaminckx et al. 1997). Such a dual model for APC may also explain why mutations that result in the complete loss of the APC protein are usually not found in human cancer cells, as loss of the entire APC protein might inhibit rather than activate β -Catenin function and may not lead to colon cancer.

Materials and methods

General methods and strains

C. elegans strains were cultured at 20°C as described in Brenner (1974). Wild type refers to *C. elegans* variety Bristol, strain N2. Pn.p cell lineages were determined by direct observation of the cell divisions using Nomarski optics, as described previously (Sternberg and Horvitz 1986). Unless noted otherwise, the mutations used have been described (Riddle et al. 1997) and are listed below.

LGI, *unc-13(e51)*, *apr-1(zh10)* (see below), *lin-10(e1439)*, *unc-29(e1072)*, *lin-11(n566)*; LGIII, *unc-119(e2498)*, *lin-39(n1880)* (Clark et al. 1993); LGIV, *him-5(e1490)*, *swls1[ceh-13::GFP]* (Wittmann et al. 1997), *gals44[lin-31::apr-1i]* (see below), *jcl1[jam-1::GFP]* (Mohler et al. 1998); LGX, *bar-1(ga80)* (Eisenmann et al. 1998); LG not determined, *zhls2[apr-1::GFP]* (see below).

Isolation of the apr-1(zh10) deletion

A library consisting of $\sim 4.8 \times 10^5$ mutagenized F₁ animals was screened for deletions in the *apr-1* locus using the strategy described (Jansen et al. 1997) except for the following modifications: Wild-type animals were treated with 75 mM EMS, allowed to recover for 16 hr and five P₀ animals were plated on each of 960 NGM plates (6-cm diam.) such that each plate contained on the average 500 mutagenized F₁ animals. The cultures were grown for 7 days before harvesting one-third of the animals for DNA isolation. To identify the plate containing the *apr-1(zh10)* deletion, DNA extracted from the individual cultures was combined in pools of 12 cultures and 50 ng of DNA from each pool was subjected to PCR analysis with the primers O96 (5'-CGTAAACGTAGGGACACCGAGGAG-

GAGG-3') and O100 (5'-CAATACGACAGTTCGACGAGC-3') for the first amplification step followed by a second amplification with the nested primers O95 (5'-CGACATGACGTCG-GTATAGGACGGCTTAG-3') and O101 (5'-AGAAGATGTT-GAAGACT-3'). The remaining animals from the culture containing the *zh10* deletion were subjected to three rounds of selection until a single animal of the genotype *apr-1(zh10)/+* had been identified. Before phenotypic analysis, the *apr-1(zh10)* mutation was backcrossed at least 10 times against *unc-29(e1072)*; *him-5(e1490)* animals.

Strain construction

Standard methods were used to construct double and triple mutants (Brenner 1974). The *apr-1(zh10)* deletion was *cis*-linked to the *unc-29(e1072)* mutation and followed by PCR amplification with the primers O95 and O101 (see above) or by observing the closely linked *Unc* phenotype. Strains heterozygous for the *apr-1(zh10)* mutation were balanced either with *lin-10(e1439)* or with *unc-13(e51) lin-11(n566)* as indicated in the Figure legends 1–4. *apr-1(zh10) unc-29(e1072)/lin-10(e1439)* animals segregated <1% viable *Unc* animals ($n > 500$) that were most likely recombinants between *apr-1* and *unc-29*, because they were heterozygous for the *apr-1(zh10)* mutation as determined by PCR analysis. Strains that were homozygous for *apr-1(zh10)* contained a rescuing *apr-1* transgene on the extrachromosomal array *zhEx11* (see below).

Molecular biology

The rescuing wild-type *apr-1* construct (*apr-1wt*) was obtained by ligation of a 7.3-kb-long PCR product amplified from wild-type genomic DNA with the primers 5'-CGAAGCCG-CAAATTCTGCGAA-3' TC and 5'-TTTCTCCCATTCCTGCGATTTC-3' into the vector pGEM-T (Invitrogen). To construct the *apr-1::GFP* reporter, a 4.8-kb *SalI*–*Bam*HI fragment isolated from *apr-1wt* was cloned into the *SalI* and *Bam*HI sites of the GFP vector pPD95.67 (gift of A. Fire, Carnegie Institution, Baltimore, MD). The plasmid *loxP* > *apr-1* > *loxP* was obtained by inserting synthetic double-stranded oligonucleotides containing the *loxP* site 5'-ATAACTTCGTATAATGTATGCTATACGAAGTTAT-3' (Kilby et al. 1993) into the *Bpu*AI and *Sph*I sites of plasmid *apr-1wt*. A clone carrying the two *loxP* sites in a parallel orientation was identified by sequence analysis. For the construction of the *lin-31::cre* plasmid, a 1.1-kb cDNA fragment containing the entire *cre* recombinase ORF was isolated by PCR amplification with the primers 5'-TTTCTC-GAGATGTCCAATTTACTGACCGTAC-3' and 5'-TTTGCG-GCCGCTAATCGCCATCTTCCAGCAGG-3' from the plasmid pCAGGS-*cre* (Araki et al. 1995), digested with *Xho*I and *Not*I and cloned into the *SalI* and *Not*I sites of plasmid pB253 (Tan et al. 1998). The plasmid *lin-31::apr-1i* was obtained by cloning a 1.3-kb *Xho*I fragment of the *apr-1* cDNA (nucleotides 85–1445 the coding sequence) into the *SalI* site of pB253 in the antisense orientation.

Antibody production, immunofluorescence, and microscopy

A 1.3-kb *apr-1* cDNA fragment was isolated from a partial cDNA clone by PCR amplification with the primers 5'-TTTG-GATCCACGGAGGATCAAGCGCAAATG-3' and 5'-TTTG-GATCCGAACAGGCAGTCTTGACTTTCC-3' and cloned into the *Bam*HI site of the *Escherichia coli* expression vector pDS6 (Stueber and Bujard 1982) to produce an APR-1 fragment consisting of amino acids 551–983 with an amino-terminal His₆ tag. The His₆::APR-1 fusion protein was produced in the *E. coli*

strain M15, purified on NTA agarose according to the manufacturer's instructions (Qiagen), and used to immunize rats and rabbits. Antisera were affinity purified against the recombinant APR-1 fragment immobilized on PVDF membranes (Immobilon).

Staining of embryos and staged larvae with rat anti-APR-1 antibodies was carried out as described (Miller and Shakes 1995), except that fixation was done in ice-cold methanol/acetone (1:1) for 5 min followed by air-drying. Rabbit anti-LIN-39 antibodies were preadsorbed against a total extract from *lin-39(n1880)* animals before use, and LIN-39 staining was performed as described (Maloof and Kenyon 1998). CEH-13::GFP was observed in living embryos mounted on agarose pads in M9 solution. In all staining experiments, the mAb MH27 (Francis and Waterston 1985) was included to label the adherens junctions. As secondary reagents, fluorescein-conjugated donkey anti-rat (Jackson laboratories), Alexa488-conjugated goat anti-rabbit (Molecular probes), and rhodamine-conjugated donkey anti-mouse antibodies (Jackson laboratories) were used. DAPI was included during the incubations with secondary antibodies at a concentration of 0.5 µg/ml to label the nuclei of the cells. Fluorescent images were recorded by confocal laser scanning microscopy (Leica TC 4D or Bio-Rad MRC 600) or with a cooled CCD camera (Hamamatsu C4742-95) on a Leica DM/IR B inverted microscope. Embryos carrying the *jcIs1* array were observed by four-dimensional microscopy using two-photon excitation as described (Mohler and White 1998; Mohler et al. 1998).

Transgenic animals

Transgenic animals were generated by microinjection of DNA into the distal arms of the gonad as described (Mello et al. 1991). Where indicated, extrachromosomal arrays were integrated into the genome by γ -irradiation of L4 larvae. Animals carrying integrated arrays were backcrossed at least five times before further analysis.

To rescue the lethal and vulvaless phenotypes caused by the *apr-1(zh10)* mutation, the plasmid *apr-1wt* was injected at a concentration of 100 ng/µl along with 100 ng/µl *sur-5::GFP* plasmid pTG96 (Yochem et al. 1998) as transformation marker into *apr-1(zh10) unc-29(e1072)/lin-10(e1439)* animals yielding the extrachromosomal array *zhEx3*. Transformants were identified with a dissecting microscope under fluorescent light illumination. *apr-1(zh10) unc-29(e1072); zhEx3[apr-1wt]* animals developed into healthy adults that were sterile, presumably because *apr-1* is required for oogenesis and the multicopy *apr-1* transgene was not active in the germ line. The sterility caused by *apr-1(zh10)* could be rescued when a low concentration of a rescue construct was injected together with an excess of genomic DNA to allow germ-line expression of the transgene as described below for the *zhEx11* array. The *zhEx2* array was obtained by injecting 100 ng/µl plasmid *apr-1wt* along with 50 ng/µl plasmid pRF4, which contains the semidominant marker *rol-6(su1006)* into N₂ animals. The plasmids *apr-1::GFP*, *lin-31::cre*, and *lin-31::apr-1i* were each injected at a concentration of 100 ng/µl together with 20 ng/µl wild-type *unc-119* plasmid (Maduro and Pilgrim 1995) into *unc-119(e2498)* animals to yield the arrays *zhIs2*, *zhEx12*, and *gals44*, respectively. The plasmid *loxP* > *apr-1* > *loxP* was linearized with *Not*I and injected at a concentration of 5 ng/µl together with 200 ng/µl yeast genomic DNA and 5 ng/µl transformation marker pTG96 linearized with *Eag*I into *apr-1(zh10) unc-29(e1072)/lin-10(e1439)* animals. The resulting array *zhEx11* rescued the sterility, lethality, and vulvaless phenotypes of *apr-1(zh10)* animals. For each construct, at least three independent lines were obtained and a representative line was kept for further analysis.

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